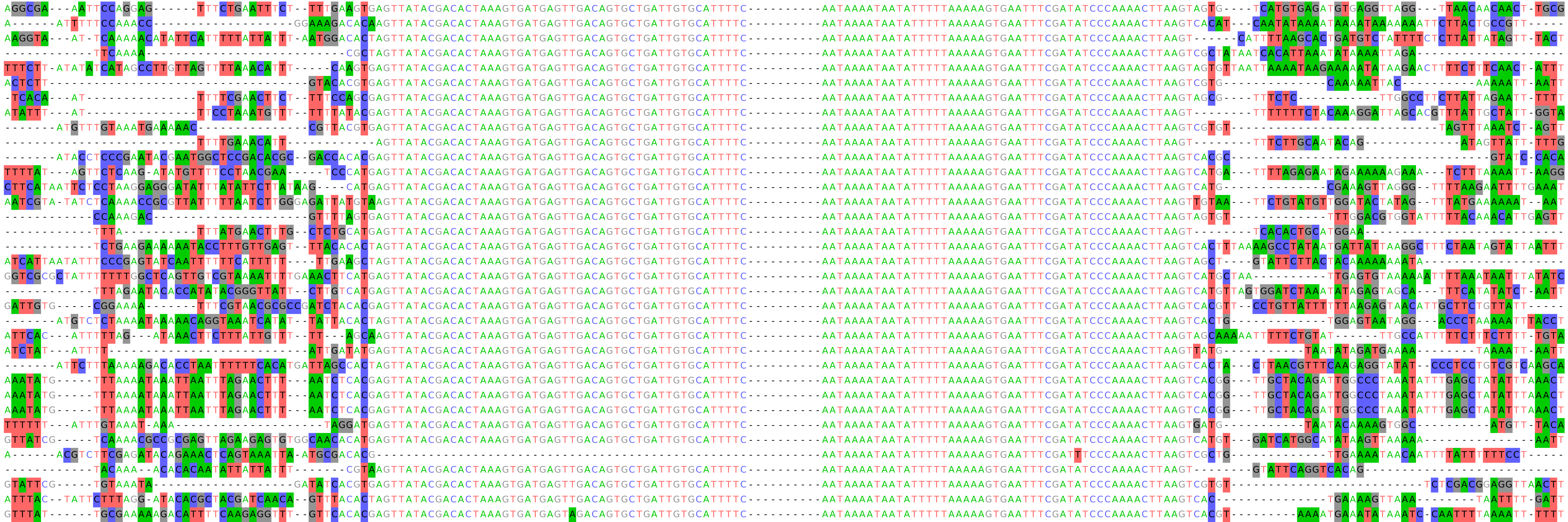
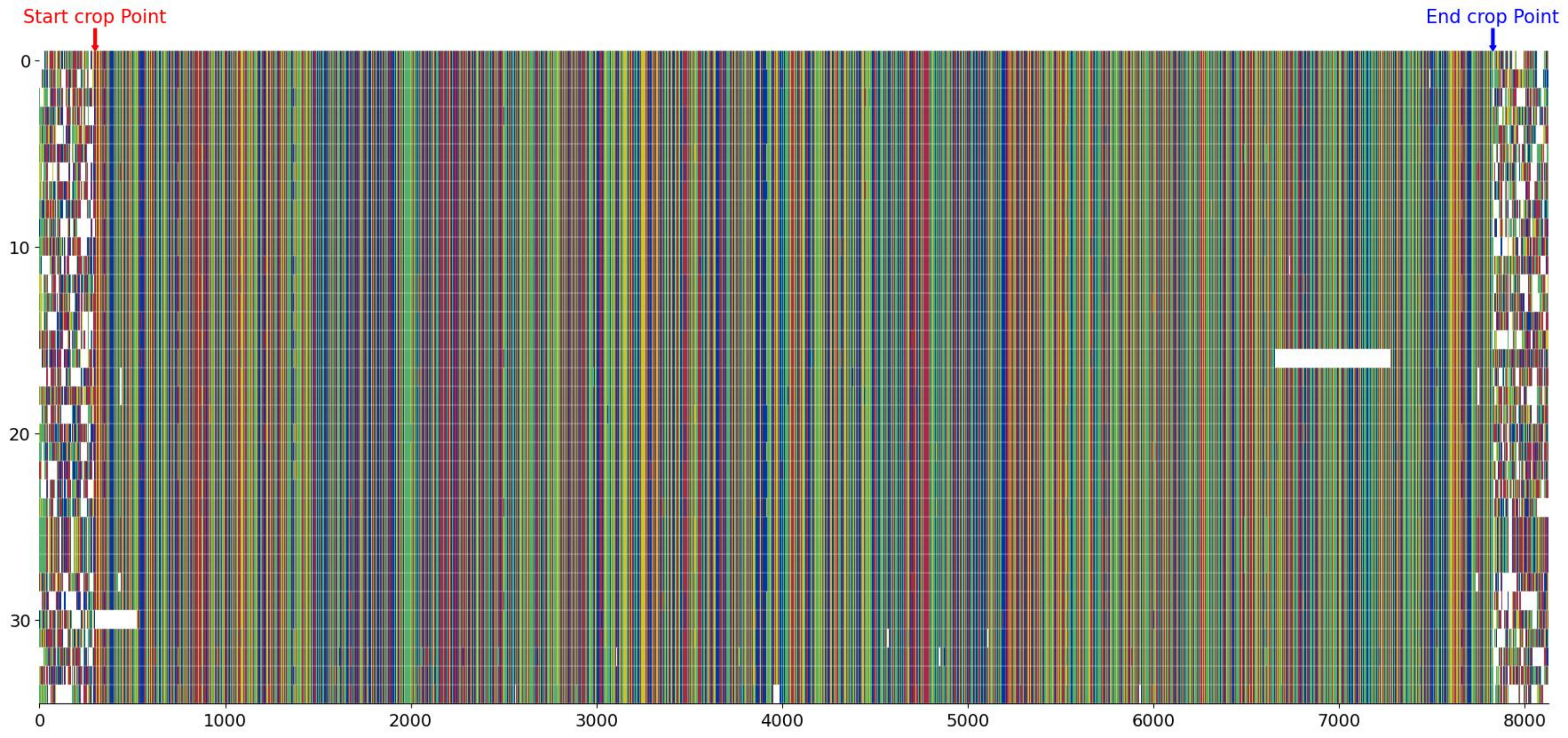


MSA length = 7531

Start crop Point

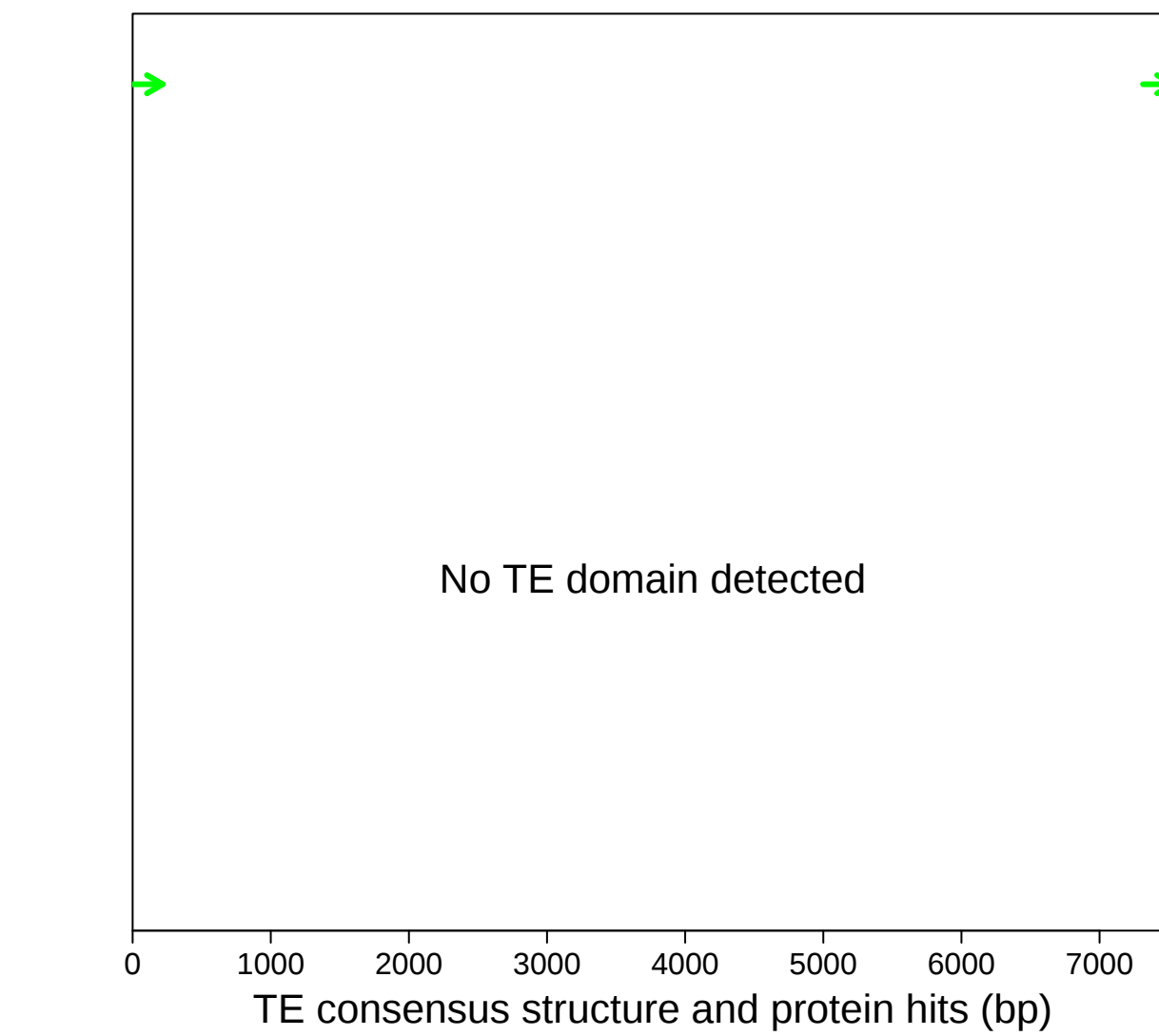
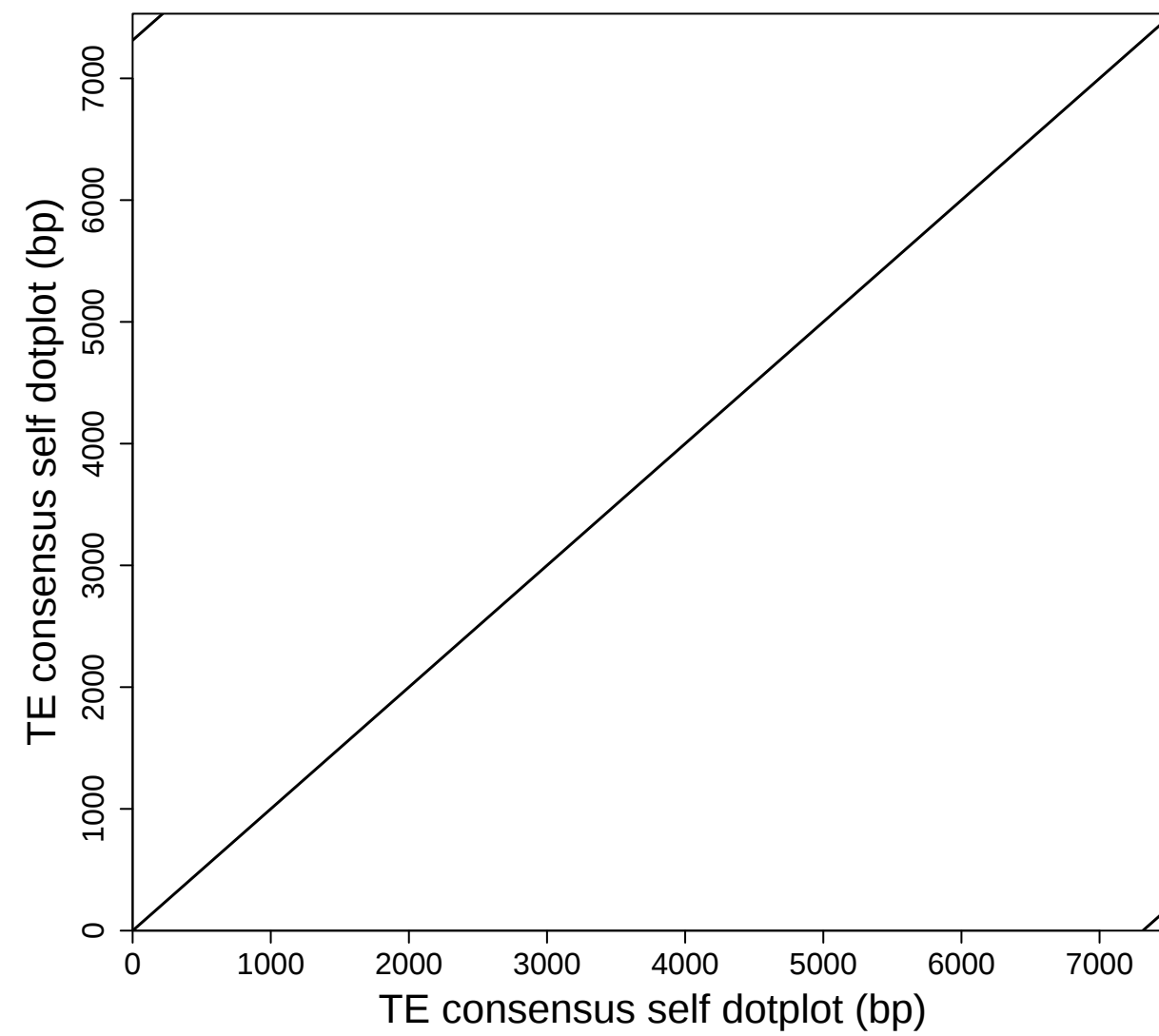
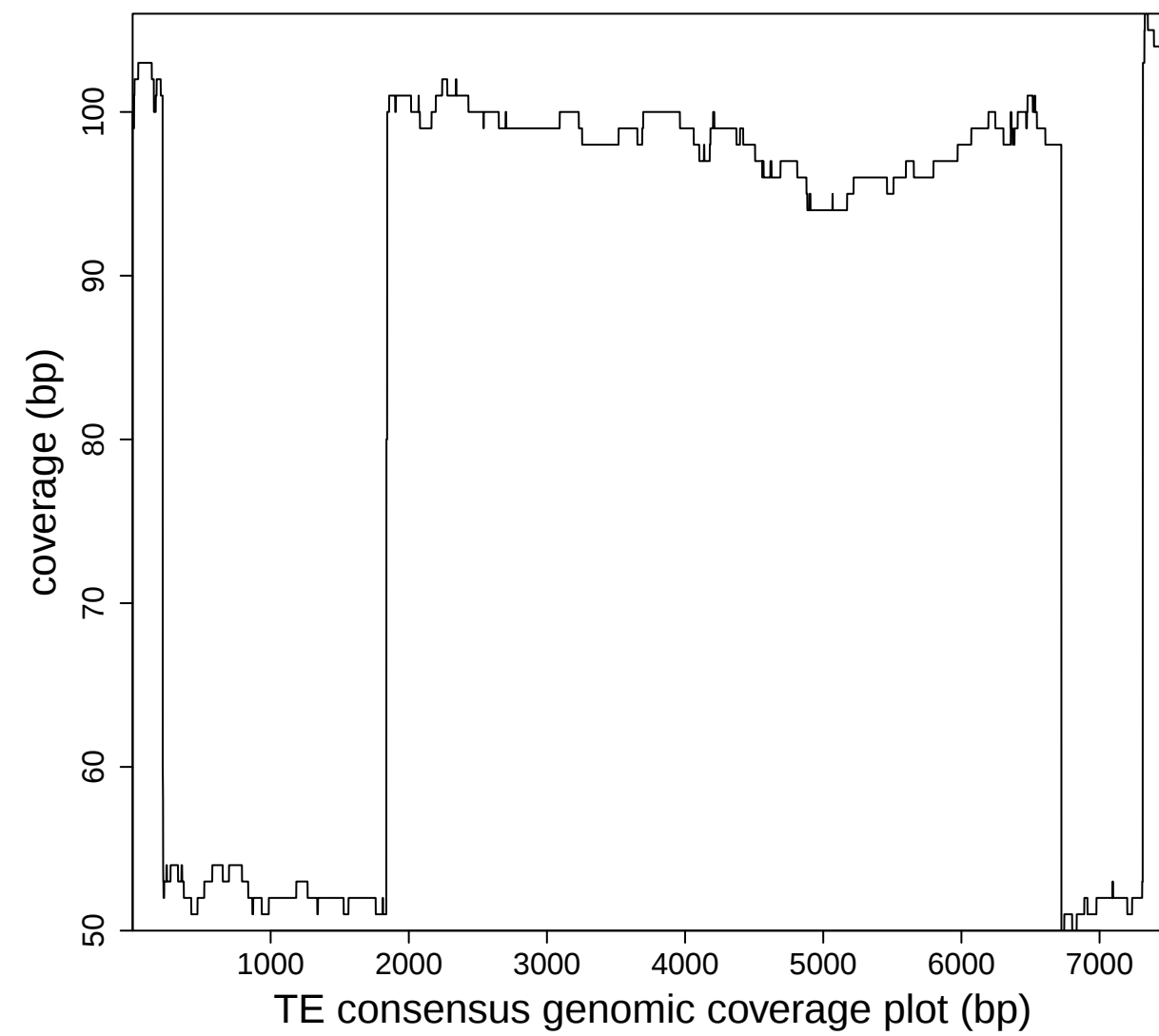
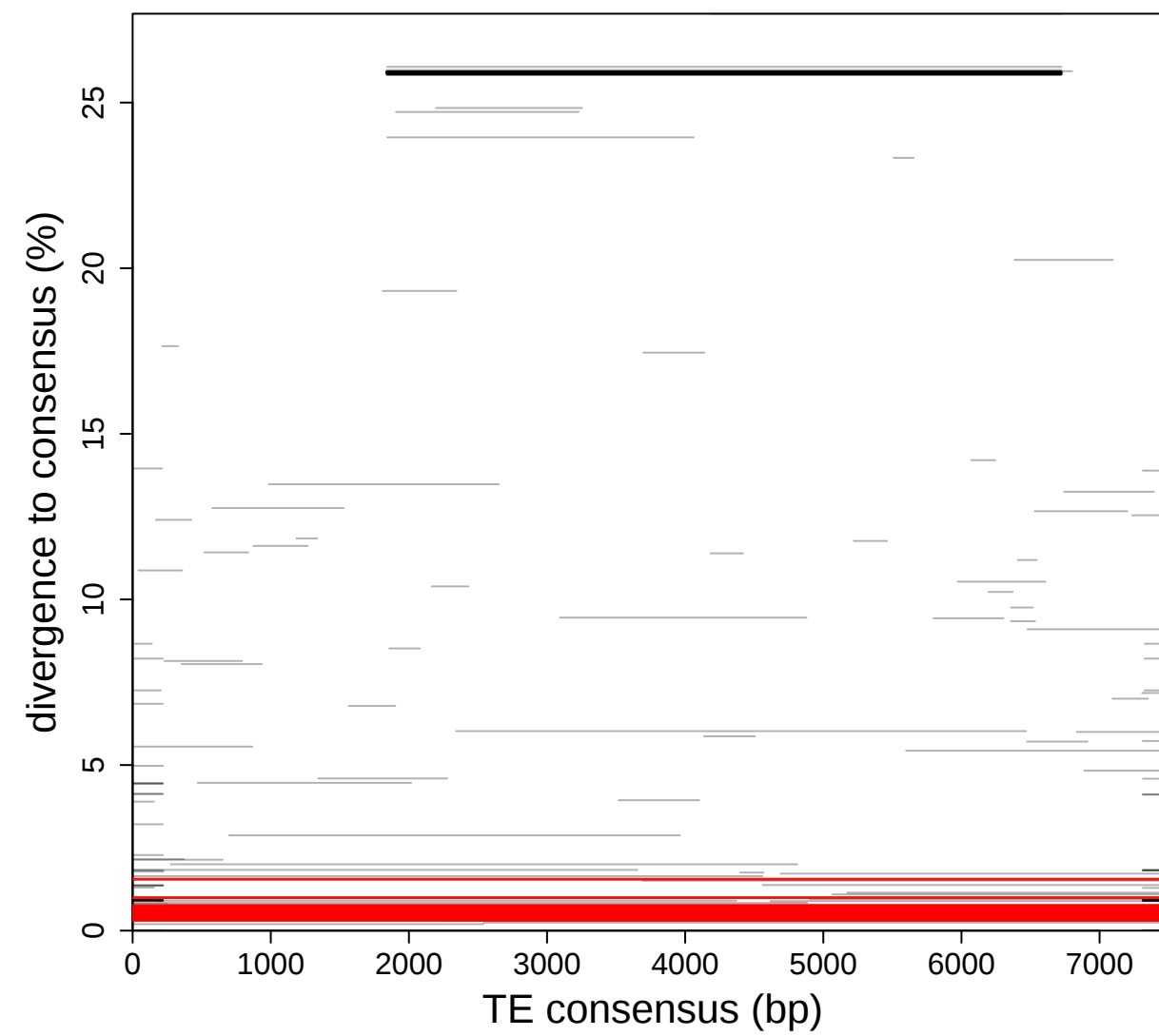
End crop Point





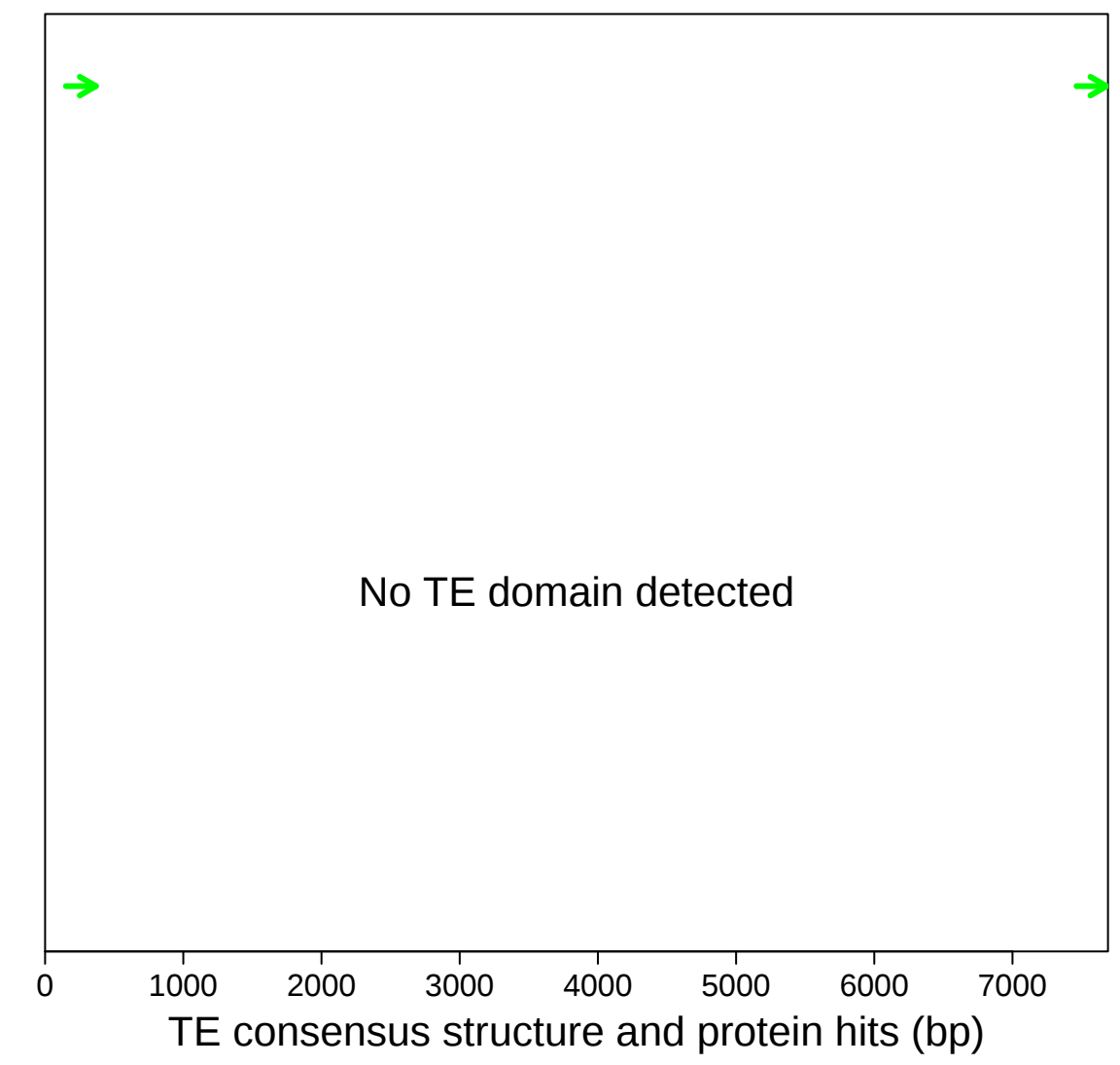
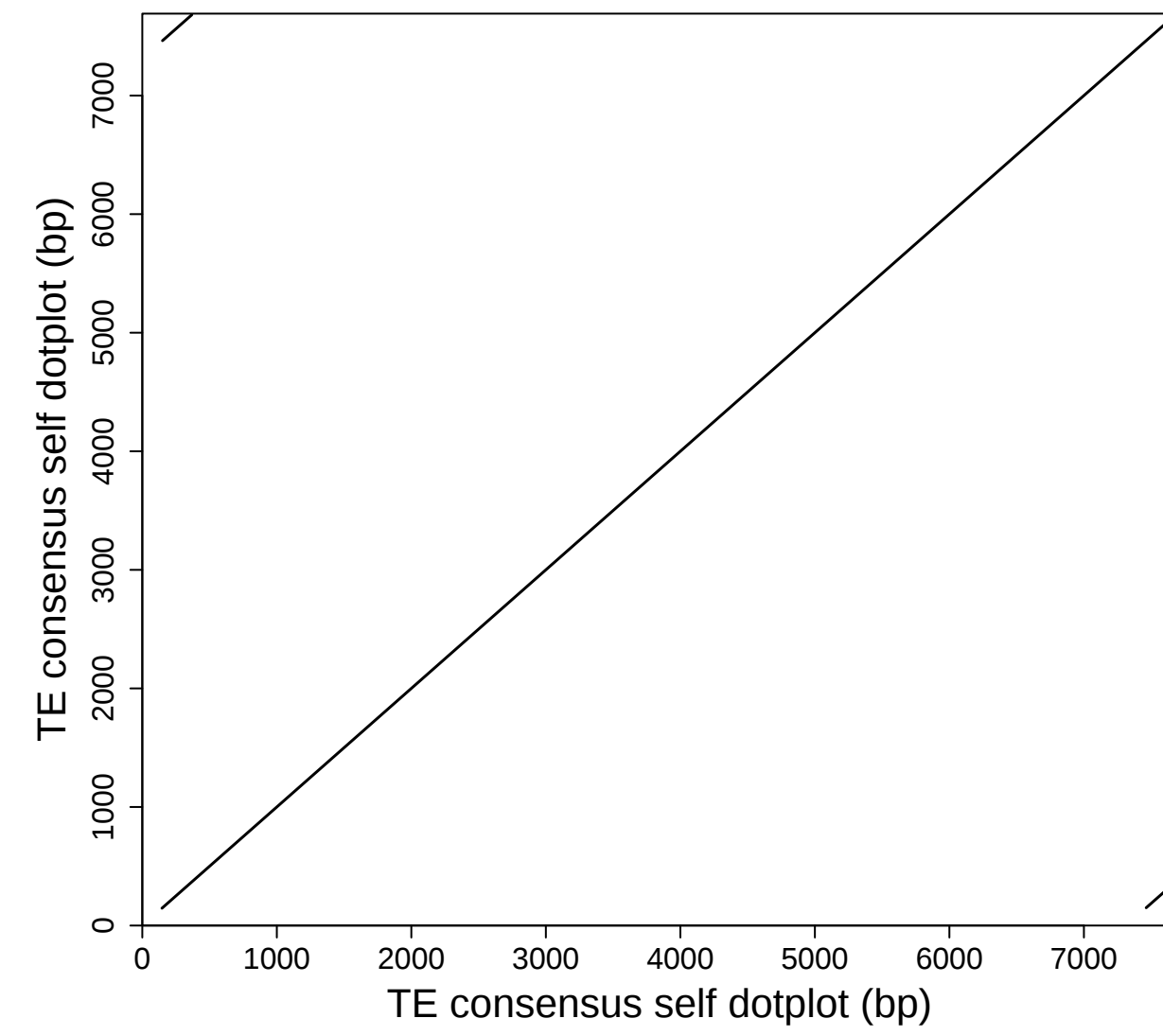
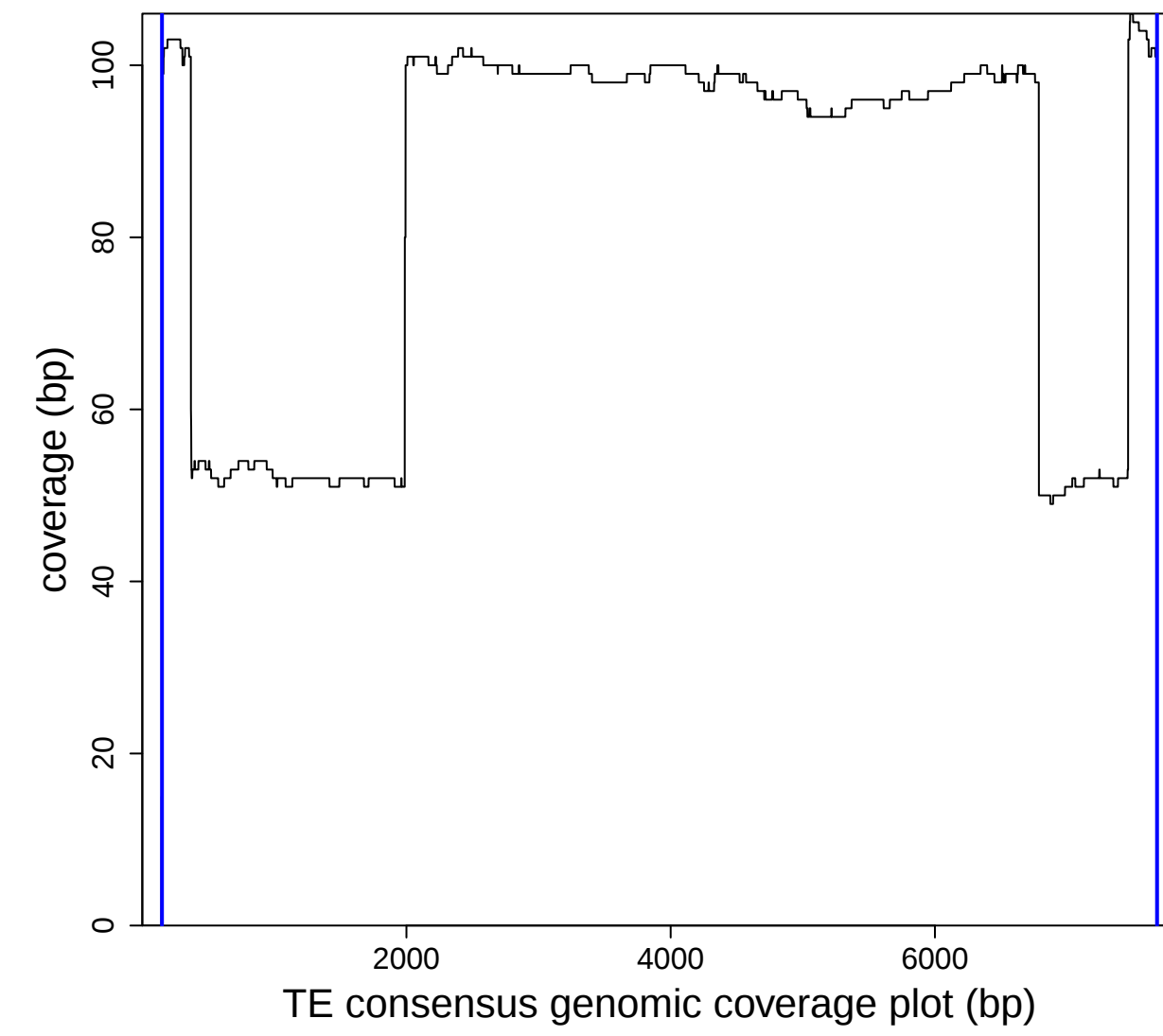
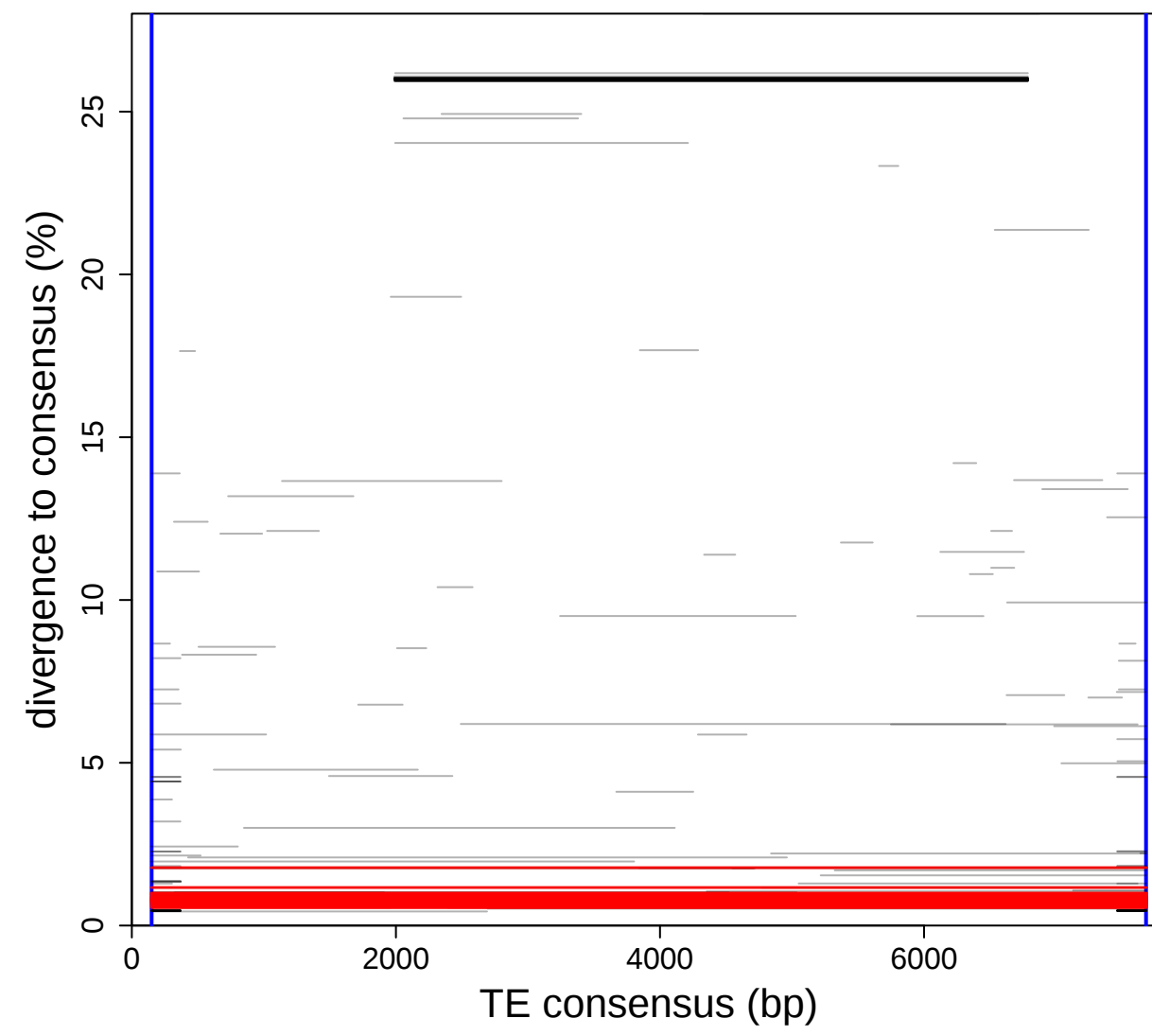
ed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.
size: 7531bp; fragments: 268; full length: 32 (≥ 6777.9 bp)

After TETrimmer 7531 bp



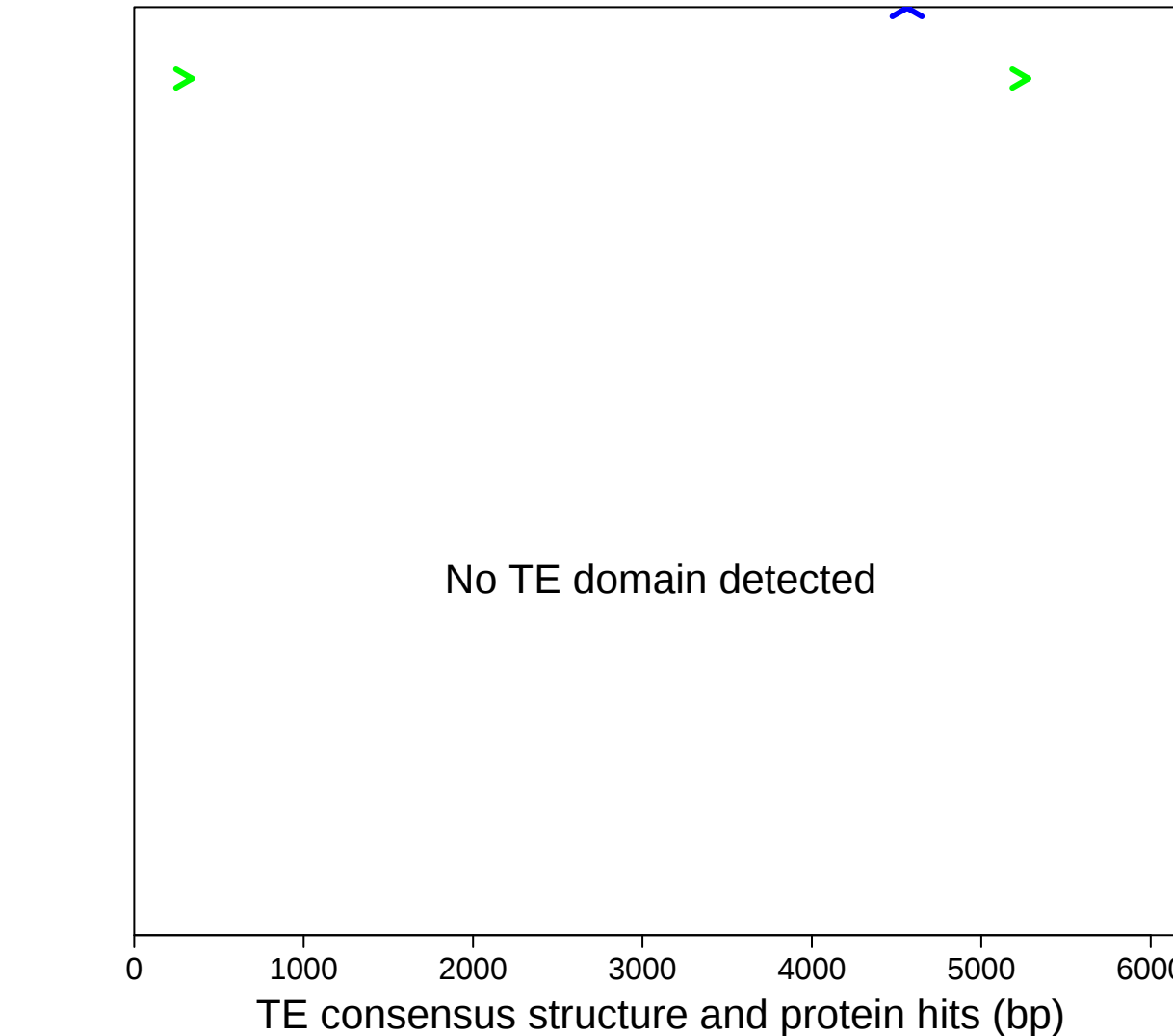
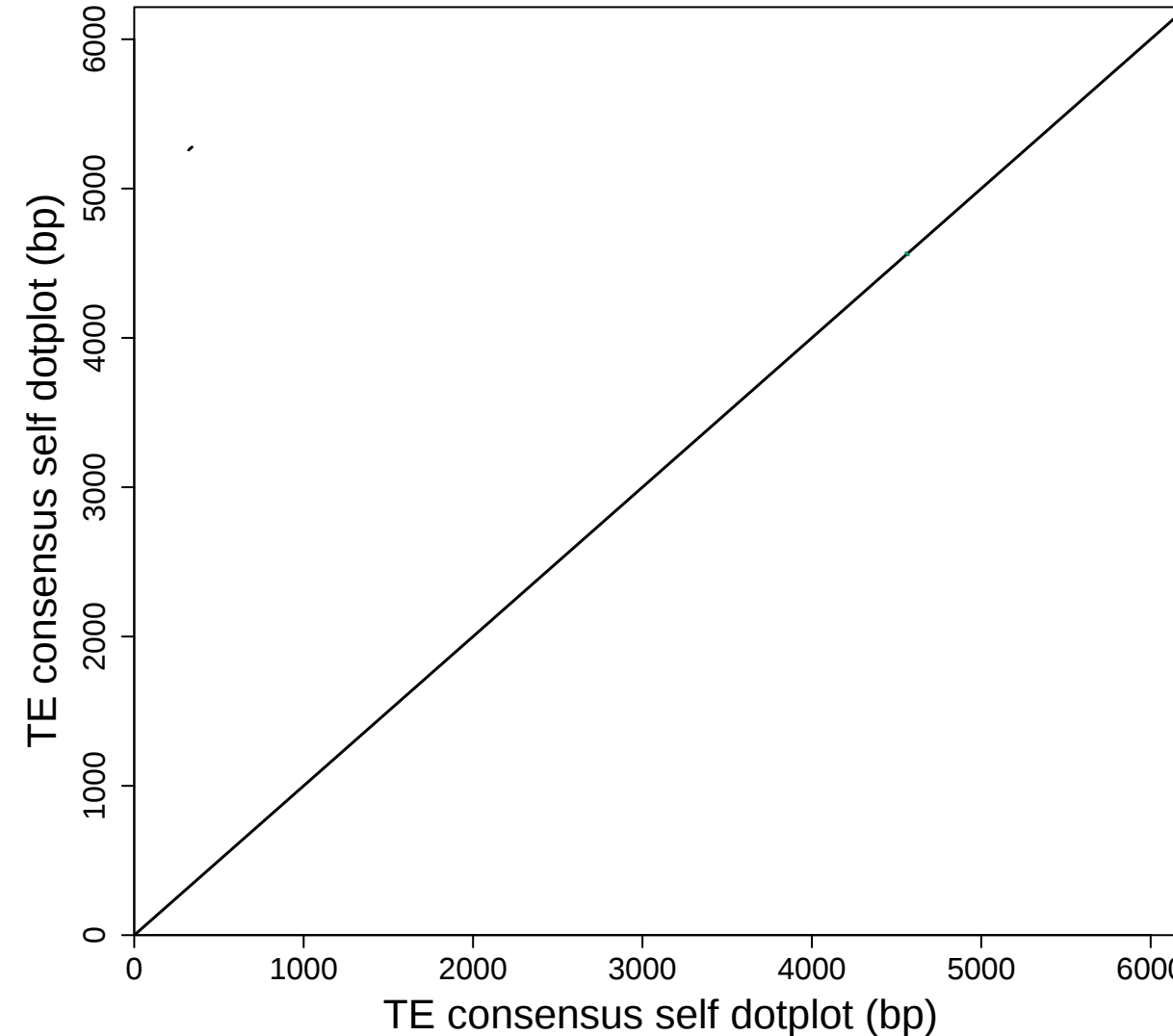
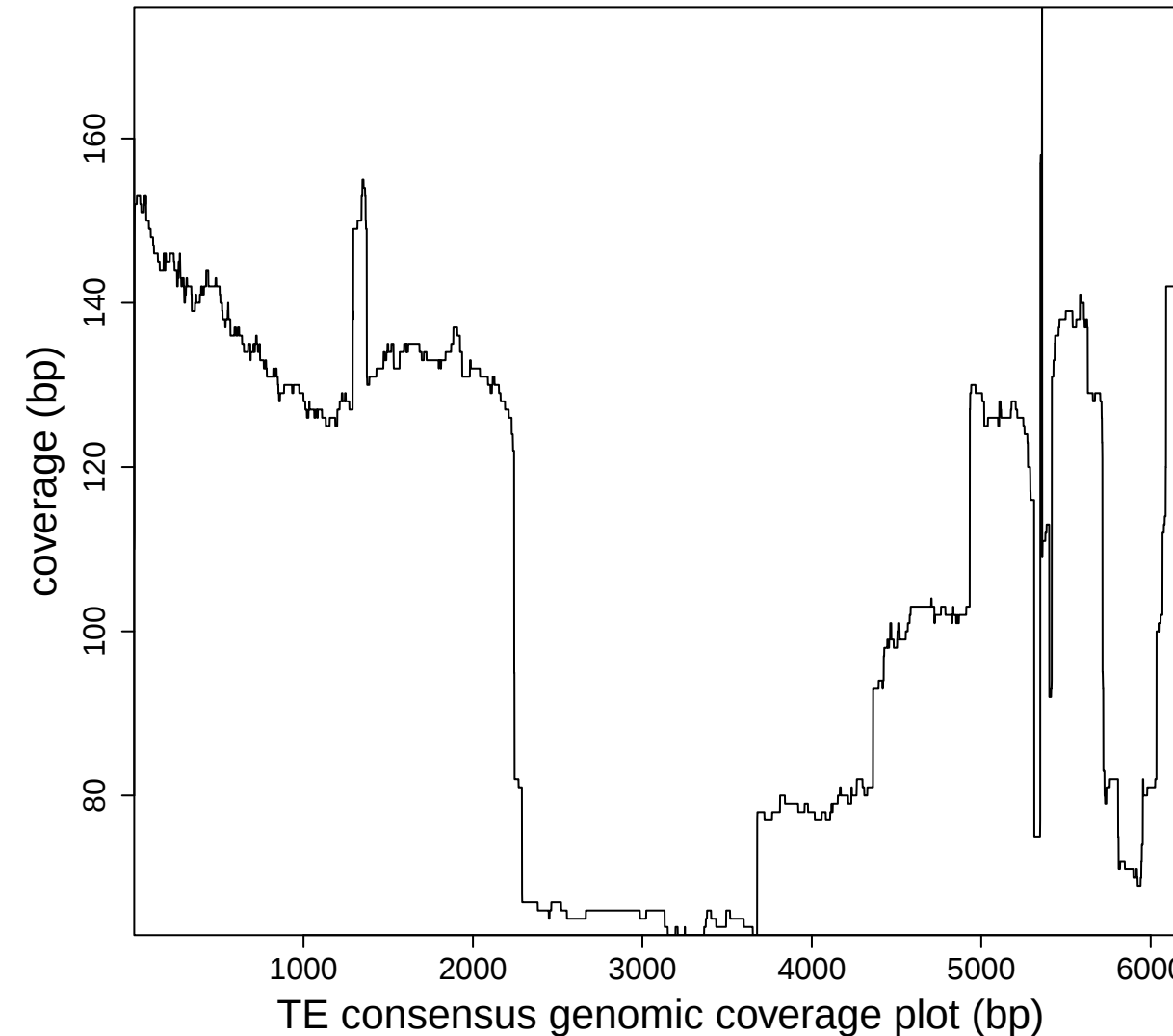
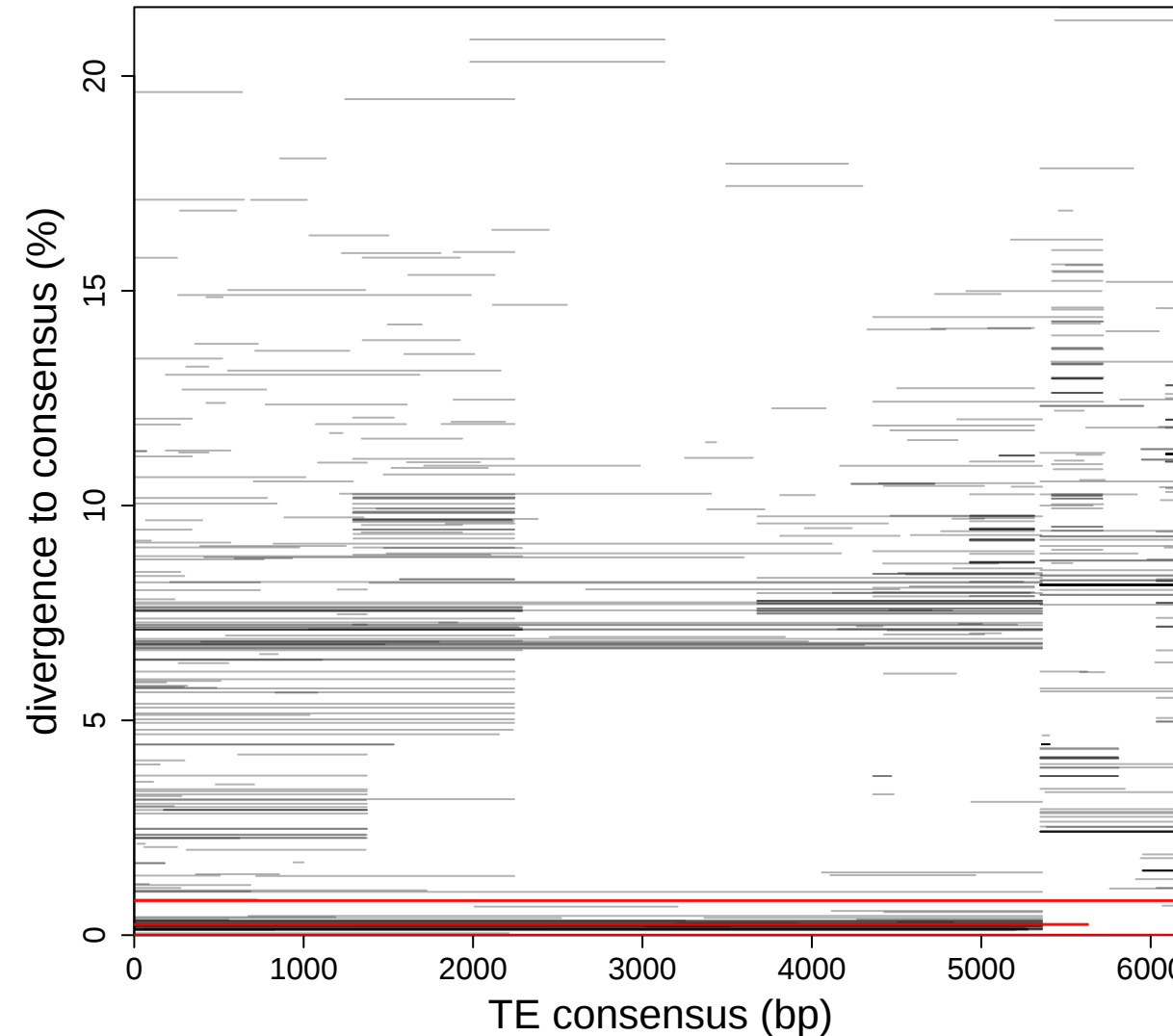
ly_696.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa
size: 7831bp; fragments: 266; full length: 32 (≥ 7047.9 bp)

After TEtrimmer Extended plot Blue lines are boundaries

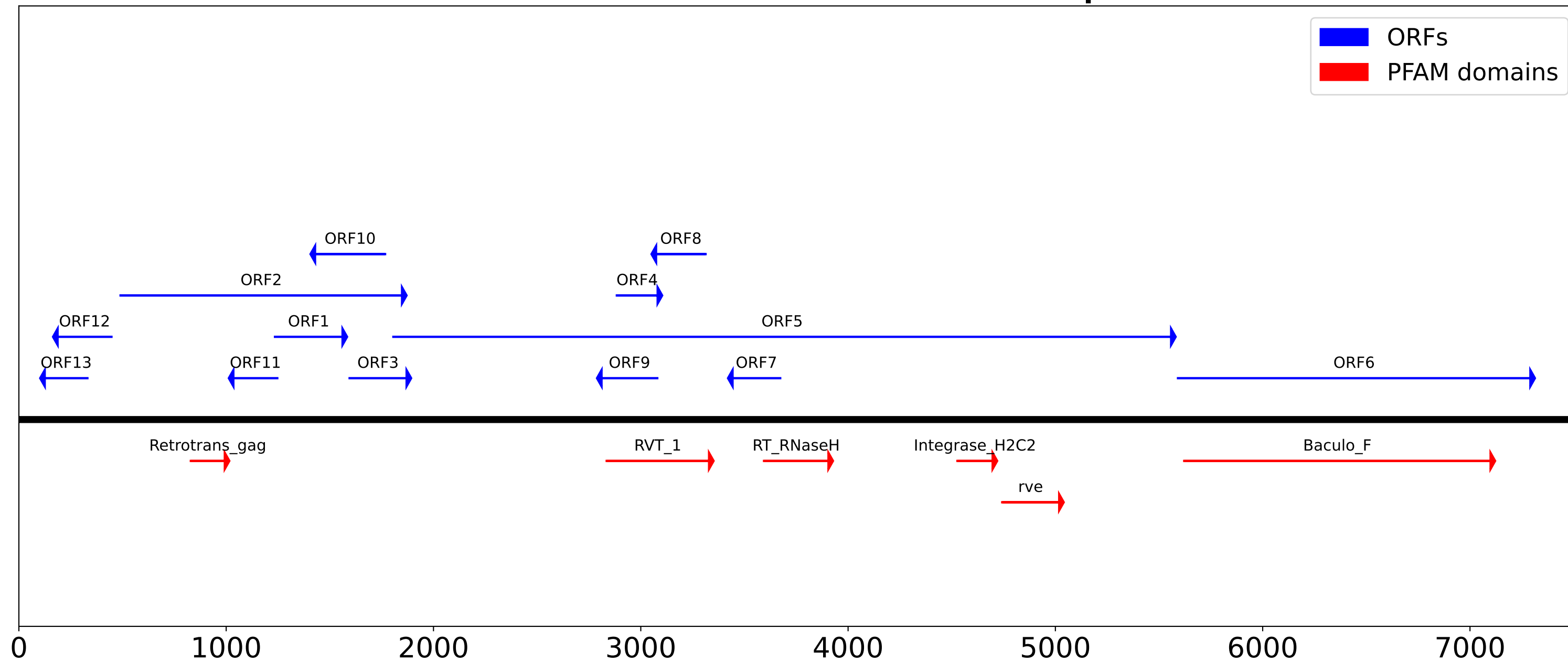


TE: ltr_1_family_696
size: 6215bp; fragments: 608; full length: 3 (>=5593.5bp)

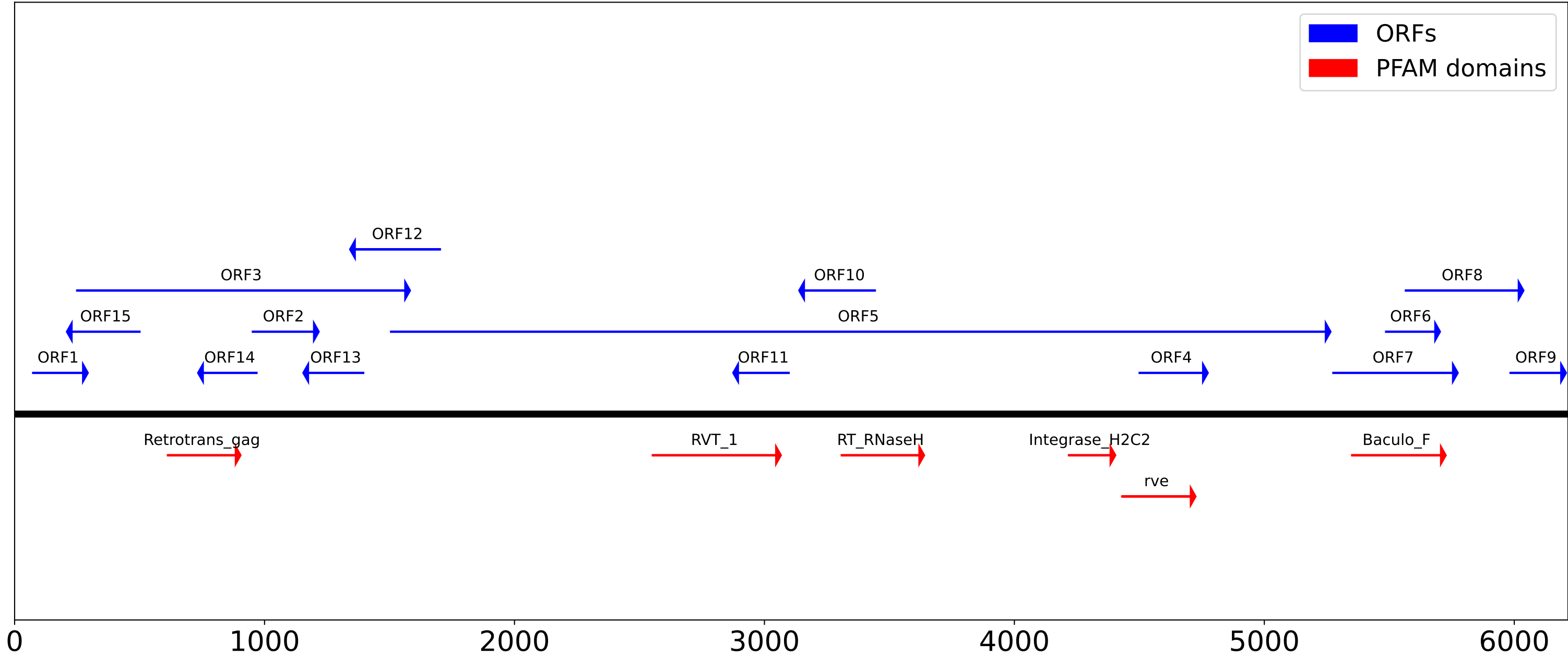
Before TEtrimmer 6215 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 09/10/25)

